

SEQUENCE LISTING

Substantially identical to sequence listing in 17282 CIP

<110> Steward, Lance E.
Aoki, K. Roger
Sachs, George

<120> Compositions, and Methods for the
Treatment of Pancreatitis

<130> 17282 CIP

<150> 09/288,326
<151> 1999-04-08

<160> 11

<170> FastSEQ for Windows Version 3.0

<210> 1
<211> 129
<212> PRT
<213> Homo sapiens

<400> 1

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Leu	Thr	Gln	Pro	Val	Pro	Pro	Ala	Asp	Pro	Ala	Gly	Ser	Gly	Leu	Gln	
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Arg	Ala	Glu	Glu	Ala	Pro	Arg	Arg	Gln	Leu	Arg	Val	Ser	Gln	Arg	Thr	
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Asp	Gly	Glu	Ser	Arg	Ala	His	Leu	Gly	Ala	Leu	Leu	Ala	Arg	Tyr	Ile	
								65			70				80	
Gln	Gln	Ala	Arg	Lys	Ala	Pro	Ser	Gly	Arg	Met	Ser	Ile	Val	Lys	Asn	
								85			90				95	
Leu	Gln	Asn	Leu	Asp	Pro	Ser	His	Arg	Ile	Ser	Asp	Arg	Asp	Tyr	Met	
								100			105				110	
Gly	Trp	Met	Asp	Phe	Gly	Arg	Arg	Ser	Ala	Glu	Glu	Tyr	Glu	Tyr	Pro	
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Ser																

<210> 2
<211> 58
<212> PRT
<213> Homo sapiens

<400> 2

Val	Ser	Gln	Arg	Thr	Asp	Gly	Glu	Ser	Arg	Ala	His	Leu	Gly	Ala	Leu
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Leu	Ala	Arg	Tyr	Ile	Gln	Gln	Ala	Arg	Lys	Ala	Pro	Ser	Gly	Arg	Met
								20			25				30
Ser	Ile	Val	Lys	Asn	Leu	Gln	Asn	Leu	Asp	Pro	Ser	His	Arg	Ile	Ser
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Asp	Arg	Asp	Tyr	Met	Gly	Trp	Met	Asp	Phe						

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55

<210> 3
<211> 39
<212> PRT
<213> Homo sapiens

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Lys Asn Leu Gln Asn Leu Asp Pro Ser His Arg Ile Ser Asp Arg Asp
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Tyr Met Gly Trp Met Asp Phe
35

<210> 4
<211> 33
<212> PRT
<213> Homo sapiens

<400> 4
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Asp Pro Ser His Arg Ile Ser Asp Arg Asp Tyr Met Gly Trp Met Asp
20 25 30
Phe

<210> 5
<211> 12
<212> PRT
<213> Homo sapiens

<400> 5
Ile Ser Asp Arg Asp Tyr Met Gly Trp Met Asp Phe
1 5 10

<210> 6
<211> 9
<212> PRT
<213> Homo sapiens

<400> 6
Arg Asp Tyr Met Gly Trp Met Asp Phe
1 5

<210> 7
<211> 448
<212> PRT
<213> Clostridium botulinum

<400> 7
Met Pro Phe Val Asn Lys Gln Phe Asn Tyr Lys Asp Pro Val Asn Gly
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Val Asp Ile Ala Tyr Ile Lys Ile Pro Asn Ala Gly Gln Met Gln Pro
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Val Lys Ala Phe Lys Ile His Asn Lys Ile Trp Val Ile Pro Glu Arg

35	40	45
Asp Thr Phe Thr Asn Pro Glu	Glu Gly Asp	Leu Asn Pro Pro Pro Glu
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Ala Lys Gln Val Pro Val Ser Tyr	Tyr Asp Ser Thr	Tyr Leu Ser Thr
65	70	75
Asp Asn Glu Lys Asp Asn Tyr	Leu Lys Gly Val	Thr Lys Leu Phe Glu
85	90	95
Arg Ile Tyr Ser Thr Asp Leu	Gly Arg Met	Leu Leu Thr Ser Ile Val
100	105	110
Arg Gly Ile Pro Phe Trp Gly	Gly Ser Thr Ile Asp	Thr Glu Leu Lys
115	120	125
Val Ile Asp Thr Asn Cys	Ile Asn Val Ile Gln	Pro Asp Gly Ser Tyr
130	135	140
Arg Ser Glu Glu Leu Asn	Leu Val Ile Ile	Gly Pro Ser Ala Asp Ile
145	150	155
Ile Gln Phe Glu Cys Lys	Ser Phe Gly His	Glu Val Leu Asn Leu Thr
165	170	175
Arg Asn Gly Tyr Gly Ser Thr	Gln Tyr Ile Arg Phe Ser	Pro Asp Phe
180	185	190
Thr Phe Gly Phe Glu Ser	Leu Glu Val Asp	Thr Asn Pro Leu Leu
195	200	205
Gly Ala Gly Lys Phe Ala	Thr Asp Pro Ala Val	Thr Leu Ala His Glu
210	215	220
Leu Ile His Ala Gly His	Arg Leu Tyr Gly	Ile Ala Ile Asn Pro Asn
225	230	235
Arg Val Phe Lys Val Asn	Thr Asn Ala Tyr	Tyr Glu Met Ser Gly Leu
245	250	255
Glu Val Ser Phe Glu Glu	Leu Arg Thr Phe Gly	Gly His Asp Ala Lys
260	265	270
Phe Ile Asp Ser Leu Gln	Glu Asn Glu Phe	Arg Leu Tyr Tyr Tyr Asn
275	280	285
Lys Phe Lys Asp Ile Ala	Ser Thr Leu Asn	Lys Ala Lys Ser Ile Val
290	295	300
Gly Thr Thr Ala Ser Leu	Gln Tyr Met Lys	Asn Val Phe Lys Glu Lys
305	310	315
Tyr Leu Leu Ser Glu Asp	Thr Ser Gly Lys	Phe Ser Val Asp Lys Leu
325	330	335
Lys Phe Asp Lys Leu	Tyr Lys Met	Leu Thr Glu Ile Tyr Thr Glu Asp
340	345	350
Asn Phe Val Lys Phe	Phe Lys Val	Leu Asn Arg Lys Thr Tyr Leu Asn
355	360	365
Phe Asp Lys Ala Val	Phe Lys Ile Asn	Ile Val Pro Lys Val Asn Tyr
370	375	380
Thr Ile Tyr Asp Gly	Phe Asn Leu Arg	Asn Thr Asn Leu Ala Ala Asn
385	390	395
Phe Asn Gly Gln Asn	Thr Glu Ile Asn	Asn Met Asn Phe Thr Lys Leu
405	410	415
Lys Asn Phe Thr Gly	Leu Phe Glu	Phe Tyr Lys Leu Leu Cys Val Arg
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Gly Ile Ile Thr Ser Lys	Thr Lys Ser	Leu Asp Lys Gly Tyr Asn Lys
435	440	445

<210> 8

<211> 423

<212> PRT

<213> Clostridium botulinum

<400> 8

Ala Leu Asn Asp Leu Cys Ile Lys Val Asn Asn Trp Asp Leu Phe Phe
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 20 25 30
 Ile Thr Ser Asp Thr Asn Ile Glu Ala Ala Glu Glu Asn Ile Ser Leu
 35 40 45
 Asp Leu Ile Gln Gln Tyr Tyr Leu Thr Phe Asn Phe Asp Asn Glu Pro
 50 55 60
 Glu Asn Ile Ser Ile Glu Asn Leu Ser Ser Asp Ile Ile Gly Gln Leu
 65 70 75 80
 Glu Leu Met Pro Asn Ile Glu Arg Phe Pro Asn Gly Lys Lys Tyr Glu
 85 90 95
 Leu Asp Lys Tyr Thr Met Phe His Tyr Leu Arg Ala Gln Glu Phe Glu
 100 105 110
 His Gly Lys Ser Arg Ile Ala Leu Thr Asn Ser Val Asn Glu Ala Leu
 115 120 125
 Leu Asn Pro Ser Arg Val Tyr Thr Phe Phe Ser Ser Asp Tyr Val Lys
 130 135 140
 Lys Val Asn Lys Ala Thr Glu Ala Ala Met Phe Leu Gly Trp Val Glu
 145 150 155 160
 Gln Leu Val Tyr Asp Phe Thr Asp Glu Thr Ser Glu Val Ser Thr Thr
 165 170 175
 Asp Lys Ile Ala Asp Ile Thr Ile Ile Pro Tyr Ile Gly Pro Ala
 180 185 190
 Leu Asn Ile Gly Asn Met Leu Tyr Lys Asp Asp Phe Val Gly Ala Leu
 195 200 205
 Ile Phe Ser Gly Ala Val Ile Leu Leu Glu Phe Ile Pro Glu Ile Ala
 210 215 220
 Ile Pro Val Leu Gly Thr Phe Ala Leu Val Ser Tyr Ile Ala Asn Lys
 225 230 235 240
 Val Leu Thr Val Gln Thr Ile Asp Asn Ala Leu Ser Lys Arg Asn Glu
 245 250 255
 Lys Trp Asp Glu Val Tyr Lys Tyr Ile Val Thr Asn Trp Leu Ala Lys
 260 265 270
 Val Asn Thr Gln Ile Asp Leu Ile Arg Lys Lys Met Lys Glu Ala Leu
 275 280 285
 Glu Asn Gln Ala Glu Ala Thr Lys Ala Ile Ile Asn Tyr Gln Tyr Asn
 290 295 300
 Gln Tyr Thr Glu Glu Glu Lys Asn Asn Ile Asn Phe Asn Ile Asp Asp
 305 310 315 320
 Leu Ser Ser Lys Leu Asn Glu Ser Ile Asn Lys Ala Met Ile Asn Ile
 325 330 335
 Asn Lys Phe Leu Asn Gln Cys Ser Val Ser Tyr Leu Met Asn Ser Met
 340 345 350
 Ile Pro Tyr Gly Val Lys Arg Leu Glu Asp Phe Asp Ala Ser Leu Lys
 355 360 365
 Asp Ala Leu Leu Lys Tyr Ile Tyr Asp Asn Arg Gly Thr Leu Ile Gly
 370 375 380
 Gln Val Asp Arg Leu Lys Asp Lys Val Asn Asn Thr Leu Ser Thr Asp
 385 390 395 400
 Ile Pro Phe Gln Leu Ser Lys Tyr Val Asp Asn Gln Arg Leu Leu Ser
 405 410 415
 Thr Phe Thr Glu Tyr Ile Lys
 420

<211> 382

<212> PRT

<213> Clostridium botulinum

<400> 9

Gln Leu Phe Asn Leu Glu Ser Ser Lys Ile Glu Val Ile Leu Lys Asn
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Ala Ile Val Tyr Asn Ser Met Tyr Glu Asn Phe Ser Thr Ser Phe Trp
20 25 30
Ile Arg Ile Pro Lys Tyr Phe Asn Ser Ile Ser Leu Asn Asn Glu Tyr
35 40 45
Thr Ile Ile Asn Cys Met Glu Asn Asn Ser Gly Trp Lys Val Ser Leu
50 55 60
Asn Tyr Gly Glu Ile Ile Trp Thr Leu Gln Asp Thr Gln Glu Ile Lys
65 70 75 80
Gln Arg Val Val Phe Lys Tyr Ser Gln Met Ile Asn Ile Ser Asp Tyr
85 90 95
Ile Asn Arg Trp Ile Phe Val Thr Ile Thr Asn Asn Arg Leu Asn Asn
100 105 110
Ser Lys Ile Tyr Ile Asn Gly Arg Leu Ile Asp Gln Lys Pro Ile Ser
115 120 125
Asn Leu Gly Asn Ile His Ala Ser Asn Asn Ile Met Phe Lys Leu Asp
130 135 140
Gly Cys Arg Asp Thr His Arg Tyr Ile Trp Ile Lys Tyr Phe Asn Leu
145 150 155 160
Phe Asp Lys Glu Leu Asn Glu Lys Glu Ile Lys Asp Leu Tyr Asp Asn
165 170 175
Gln Ser Asn Ser Gly Ile Leu Lys Asp Phe Trp Gly Asp Tyr Leu Gln
180 185 190
Tyr Asp Lys Pro Tyr Tyr Met Leu Asn Leu Tyr Asp Pro Asn Lys Tyr
195 200 205
Val Asp Val Asn Asn Val Gly Ile Arg Gly Tyr Met Tyr Leu Lys Gly
210 215 220
Pro Arg Gly Ser Val Met Thr Thr Asn Ile Tyr Leu Asn Ser Ser Leu
225 230 235 240
Tyr Arg Gly Thr Lys Phe Ile Ile Lys Lys Tyr Ala Ser Gly Asn Lys
245 250 255
Asp Asn Ile Val Arg Asn Asn Asp Arg Val Tyr Ile Asn Val Val Val
260 265 270
Lys Asn Lys Glu Tyr Arg Leu Ala Thr Asn Ala Ser Gln Ala Gly Val
275 280 285
Glu Lys Ile Leu Ser Ala Leu Glu Ile Pro Asp Val Gly Asn Leu Ser
290 295 300
Gln Val Val Val Met Lys Ser Lys Asn Asp Gln Gly Ile Thr Asn Lys
305 310 315 320
Cys Lys Met Asn Leu Gln Asp Asn Asn Gly Asn Asp Ile Gly Phe Ile
325 330 335
Gly Phe His Gln Phe Asn Asn Ile Ala Lys Leu Val Ala Ser Asn Trp
340 345 350
Tyr Asn Arg Gln Ile Glu Arg Ser Ser Arg Thr Leu Gly Cys Ser Trp
355 360 365
Glu Phe Ile Pro Val Asp Asp Gly Trp Gly Glu Arg Pro Leu
370 375 380

<210> 10

<211> 4835

<212> DNA

<213> Clostridium botulinum

<400> 10

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tattnatttc	caattgtta	accctatctt	ataacggtaa	atatastatgt	ttatctatga	240
aagatgaaaa	ccataattgg	atgatatgta	ataatgataat	gtcaaagtat	ttgtatttat	300
ggtcatttaa	ataattaataa	attnaaatta	ttttaataat	tataagaggt	gtttaataatg	360
ccatttgtt	ataaaacaatt	taattataaa	gatcctgtaa	atgggtgtta	tattgcttat	420
ataaaaattc	caaatgcagg	acaaatgcaa	ccagtaaaag	cttttaataat	tcataataaa	480
atatgggtt	ttccagaaag	agatacattt	acaaatcctg	aagaaggaga	tttaataatcca	540
ccaccagaag	caaacaagaat	tccagttca	tattatgatt	caacatattt	aagtacagat	600
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acaatagata	cagaattaaa	agttattgtat	actaattgtt	ttaatgtat	acaaccagat	780
ggtagttata	gatcagaaga	acttaatcta	gtaataataa	gaccctcagc	tgatattata	840
cagtttgaat	gtaaaagctt	tggacatgaa	gttttgaatc	ttacgcgaaa	tggttatggc	900
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gttgatacaa	atccttttt	aggtgcaggc	aaatttgcta	cagatccagc	agtaacattt	1020
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aagtatgagt	tagataaata	tactatgtt	cattatctt	gtgctcaaga	atttgaacat	2040
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<210> 11
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 <213> Homo sapiens

<400> 11

Glu	Pro	Lys	Ser	Cys	Asp	Lys	Thr	His	Thr	Cys	Pro	Pro	Cys	Pro
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